

SEQUENCE LISTING

<110> Nakayama, Naoki
Wen, Duanzhi
Han, Chun-ya
He, Ching
Yu, Dongyin

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545					550						555					560
Leu	His	Tyr	Glu	Val	Leu	Leu	Ala	Gly	Leu	Gly	Gly	Ser	Glu	Gln	Gly	
				565						570				575		
Thr	Val	Thr	Ala	His	Leu	Leu	Gly	Pro	Pro	Gly	Met	Pro	Gly	Pro	Gln	
			580						585				590			
Arg	Leu	Leu	Lys	Gly	Phe	Tyr	Gly	Ser	Glu	Ala	Gln	Gly	Val	Val	Lys	
		595						600				605				
Asp	Leu	Glu	Pro	Val	Leu	Leu	Arg	His	Leu	Ala	Gln	Gly	Thr	Ala	Ser	
		610						615				620				
Leu	Leu	Ile	Thr	Thr	Lys	Ser	Ser	Pro	Arg	Gly	Glu	Leu	Arg	Gly	Gln	
625					630						635				640	
Val	His	Ile	Ala	Ser	Gln	Cys	Glu	Ala	Gly	Gly	Leu	Arg	Leu	Ala	Ser	
				645						650				655		
Glu	Gly	Val	Gln	Met	Pro	Leu	Ala	Pro	Asn	Gly	Glu	Ala	Ala	Thr	Ser	
			660						665				670			
Pro	Met	Leu	Pro	Ala	Gly	Pro	Gly	Pro	Glu	Ala	Pro	Val	Pro	Ala	Lys	
		675						680				685				
His	Gly	Ser	Pro	Gly	Arg	Pro	Arg	Asp	Pro	Asn	Thr	Cys	Phe	Phe	Glu	
		690						695				700				
Gly	Gln	Gln	Arg	Pro	His	Gly	Ala	Arg	Trp	Ala	Pro	Asn	Tyr	Asp	Pro	
705					710						715				720	
Leu	Cys	Ser	Leu	Cys	Ile	Cys	Gln	Arg	Arg	Thr	Val	Ile	Cys	Asp	Pro	
				725						730				735		
Val	Val	Cys	Pro	Pro	Pro	Ser	Cys	Pro	His	Pro	Val	Gln	Ala	Leu	Asp	
			740						745				750			
Gln	Cys	Cys	Pro	Val	Cys	Pro	Glu	Lys	Gln	Arg	Ser	Arg	Asp	Leu	Pro	
		755						760				765				
Ser	Leu	Pro	Asn	Leu	Glu	Pro	Gly	Glu	Gly	Cys	Tyr	Phe	Asp	Gly	Asp	
		770						775				780				
Arg	Ser	Trp	Arg	Ala	Ala	Gly	Thr	Arg	Trp	His	Pro	Val	Val	Pro	Pro	

785		790		795		800
Phe Gly Leu Ile	Lys Cys Ala Val Cys Thr Cys Lys Gly Ala Thr Gly					
	805		810		815	
Glu Val His Cys Glu Lys Val Gln Cys Pro Arg Leu Ala Cys Ala Gln						
	820		825		830	
Pro Val Arg Ala Asn Pro Thr Asp Cys Cys Lys Gln Cys Pro Val Gly						
	835		840		845	
Ser Gly Thr Asn Ala Lys Leu Gly Asp Pro Met Gln Ala Asp Gly Pro						
	850		855		860	
Arg Gly Cys Arg Phe Ala Gly Gln Trp Phe Pro Glu Asn Gln Ser Trp						
	865		870		875	880
His Pro Ser Val Pro Pro Phe Gly Glu Met Ser Cys Ile Thr Cys Arg						
	885		890		895	
Cys Gly Ala Gly Val Pro His Cys Glu Arg Asp Asp Cys Ser Pro Pro						
	900		905		910	
Leu Ser Cys Gly Ser Gly Lys Glu Ser Arg Cys Cys Ser His Cys Thr						
	915		920		925	
Ala Gln Arg Ser Ser Glu Thr Arg Thr Leu Pro Glu Leu Glu Lys Glu						
	930		935		940	
Ala Glu His Ser Val Asp Gly Ser Gly Thr Asp Tyr Lys Asp Asp Asp						
	945		950		955	960
Asp Lys						

<210> 11
 <211> 431
 <212> PRT
 <213> Homo sapiens

<400> 11
 Glu Gln Val Lys His Ser Glu Thr Tyr Cys Met Phe Gln Asp Lys Lys
 1 5 10 15
 Tyr Arg Val Gly Glu Arg Trp His Pro Tyr Leu Glu Pro Tyr Gly Leu
 20 25 30
 Val Tyr Cys Val Asn Cys Ile Cys Ser Glu Asn Gly Asn Val Leu Cys
 35 40 45
 Ser Arg Val Arg Cys Pro Asn Val His Cys Leu Ser Pro Val His Ile
 50 55 60
 Pro His Leu Cys Cys Pro Arg Cys Pro Glu Asp Ser Leu Pro Pro Val
 65 70 75 80

Arg Thr Thr Leu Ser Gln Trp Lys Ile Phe Thr Glu Gly Glu Ala Gln
385 390 395 400

Ile Ser Gln Met Cys Ser Ser Arg Val Cys Arg Thr Glu Leu Glu Asp
405 410 415

Leu Val Lys Val Leu Tyr Leu Glu Arg Ser Glu Lys Gly His Cys
420 425 430

<210> 12
<211> 434
<212> PRT
<213> Homo sapiens

<400> 12
Gly Lys Thr Glu Gln Val Lys His Ser Glu Thr Tyr Cys Met Phe Gln
1 5 10 15

Asp Lys Lys Tyr Arg Val Gly Glu Arg Trp His Pro Tyr Leu Glu Pro
20 25 30

Tyr Gly Leu Val Tyr Cys Val Asn Cys Ile Cys Ser Glu Asn Gly Asn
35 40 45

Val Leu Cys Ser Arg Val Arg Cys Pro Asn Val His Cys Leu Ser Pro
50 55 60

Val His Ile Pro His Leu Cys Cys Pro Arg Cys Pro Glu Asp Ser Leu
65 70 75 80

Pro Pro Val Asn Asn Lys Val Thr Ser Lys Ser Cys Glu Tyr Asn Gly
85 90 95

Thr Thr Tyr Gln His Gly Glu Leu Phe Val Ala Glu Gly Leu Phe Gln
100 105 110

Asn Arg Gln Pro Asn Gln Cys Thr Gln Cys Ser Cys Ser Glu Gly Asn
115 120 125

Val Tyr Cys Gly Leu Lys Thr Cys Pro Lys Leu Thr Cys Ala Phe Pro
130 135 140

Val Ser Val Pro Asp Ser Cys Cys Arg Val Cys Arg Gly Asp Gly Glu
145 150 155 160

Leu Ser Trp Glu His Ser Asp Gly Asp Ile Phe Arg Gln Pro Ala Asn
165 170 175

Arg Glu Ala Arg His Ser Tyr His Arg Ser His Tyr Asp Pro Pro Pro
180 185 190

Ser Arg Gln Ala Gly Gly Leu Ser Arg Phe Pro Gly Ala Arg Ser His
195 200 205

Arg Gly Ala Leu Met Asp Ser Gln Gln Ala Ser Gly Thr Ile Val Gln
210 215 220

Ile	Val	Ile	Asn	Asn	Lys	His	Lys	His	Gly	Gln	Val	Cys	Val	Ser	Asn	
225					230					235					240	
Gly	Lys	Thr	Tyr	Ser	His	Gly	Glu	Ser	Trp	His	Pro	Asn	Leu	Arg	Ala	
				245					250					255		
Phe	Gly	Ile	Val	Glu	Cys	Val	Leu	Cys	Thr	Cys	Asn	Val	Thr	Lys	Gln	
			260					265					270			
Glu	Cys	Lys	Lys	Ile	His	Cys	Pro	Asn	Arg	Tyr	Pro	Cys	Lys	Tyr	Pro	
		275					280					285				
Gln	Lys	Ile	Asp	Gly	Lys	Cys	Cys	Lys	Val	Cys	Pro	Gly	Lys	Lys	Ala	
	290					295					300					
Lys	Glu	Glu	Leu	Pro	Gly	Gln	Ser	Phe	Asp	Asn	Lys	Gly	Tyr	Phe	Cys	
305					310					315					320	
Gly	Glu	Glu	Thr	Met	Pro	Val	Tyr	Glu	Ser	Val	Phe	Met	Glu	Asp	Gly	
				325					330					335		
Glu	Thr	Thr	Arg	Lys	Ile	Ala	Leu	Glu	Thr	Glu	Arg	Pro	Pro	Gln	Val	
			340					345					350			
Glu	Val	His	Val	Trp	Thr	Ile	Arg	Lys	Gly	Ile	Leu	Gln	His	Phe	His	
		355					360					365				
Ile	Glu	Lys	Ile	Ser	Lys	Arg	Met	Phe	Glu	Glu	Leu	Pro	His	Phe	Lys	
	370					375					380					
Leu	Val	Thr	Arg	Thr	Thr	Leu	Ser	Gln	Trp	Lys	Ile	Phe	Thr	Glu	Gly	
385					390					395					400	
Glu	Ala	Gln	Ile	Ser	Gln	Met	Cys	Ser	Ser	Arg	Val	Cys	Arg	Thr	Glu	
				405					410					415		
Leu	Glu	Asp	Leu	Val	Lys	Val	Leu	Tyr	Leu	Glu	Arg	Ser	Glu	Lys	Gly	
			420					425					430			

His Cys

<210> 13
 <211> 360
 <212> PRT
 <213> Rattus norvegicus

<400> 13
 Glu Pro Val Lys His Ser Glu Thr Tyr Cys Met Phe Gln Asp Lys Lys
 1 5 10 15
 Tyr Arg Val Gly Glu Lys Trp His Pro Tyr Leu Glu Pro Tyr Gly Leu
 20 25 30
 Val Tyr Cys Val Asn Cys Ile Cys Ser Glu Asn Gly Asn Val Leu Cys

35					40					45					
Ser	Arg	Val	Arg	Cys	Pro	Thr	Leu	His	Cys	Leu	Ser	Pro	Val	His	Ile
50						55				60					
Pro	His	Leu	Cys	Cys	Pro	Arg	Cys	Pro	Asp	Ser	Leu	Pro	Pro	Met	Asn
65				70						75				80	
Asn	Lys	Val	Thr	Ser	Lys	Ser	Cys	Glu	Tyr	Asn	Gly	Thr	Thr	Tyr	Gln
				85				90						95	
His	Gly	Glu	Leu	Phe	Ile	Ala	Glu	Gly	Leu	Phe	Gln	Asn	Arg	Gln	Pro
		100						105				110			
Asn	Gln	Cys	Ser	Gln	Cys	Ser	Cys	Ser	Glu	Gly	Asn	Val	Tyr	Cys	Gly
		115				120						125			
Leu	Lys	Thr	Cys	Pro	Lys	Leu	Thr	Cys	Ala	Phe	Pro	Val	Ser	Val	Pro
130						135				140					
Asp	Ser	Cys	Cys	Arg	Val	Cys	Arg	Gly	Asp	Gly	Glu	Leu	Ser	Trp	Glu
145				150						155				160	
His	Ser	Asp	Ala	Asp	Ile	Phe	Arg	Gln	Pro	Ala	Asn	Arg	Glu	Ala	Arg
				165				170						175	
His	Ser	Tyr	Leu	Arg	Ser	Pro	Tyr	Asp	Pro	Pro	Pro	Ser	Arg	Gln	Ala
		180						185				190			
Gly	Gly	Leu	Pro	Arg	Phe	Ala	Gly	Ser	Arg	Ser	His	Arg	Gly	Ala	Val
		195				200						205			
Ile	Asp	Ser	Gln	Gln	Ala	Ser	Gly	Thr	Ile	Val	Gln	Ile	Val	Ile	Asn
210						215				220					
Asn	Lys	His	Lys	His	Gly	Gln	Val	Cys	Val	Ser	Asn	Gly	Lys	Thr	Tyr
225				230						235				240	
Ser	His	Gly	Glu	Ser	Trp	His	Ser	Asn	Leu	Arg	Ala	Phe	Gly	Ile	Val
				245				250						255	
Glu	Cys	Val	Leu	Cys	Thr	Cys	Asn	Val	Thr	Lys	Gln	Glu	Cys	Lys	Lys
		260						265				270			
Ile	His	Cys	Pro	Asn	Arg	Tyr	Pro	Cys	Lys	Tyr	Pro	Gln	Lys	Leu	Asp
		275				280						285			
Gly	Lys	Cys	Cys	Lys	Val	Cys	Pro	Glu	Glu	Pro	Pro	Ser	Gln	Asn	Phe
290						295				300					
Asp	Ser	Lys	Gly	Ser	Phe	Cys	Gly	Glu	Glu	Thr	Met	Pro	Val	Tyr	Glu
305				310						315				320	
Ala	Val	Leu	Val	Glu	Asp	Gly	Glu	Thr	Ala	Arg	Lys	Val	Ala	Leu	Glu
				325				330						335	
Thr	Glu	Lys	Pro	Pro	Gln	Val	Val	Gly	Ser	Arg	Leu	Asp	Tyr	Ser	Lys

340

345

350

Gly His Ser Pro Ala Leu Pro His
 355 360

<210> 14
 <211> 20
 <212> DNA
 <213> Mus musculus

<400> 14
 agtgcccagc tttagtccac 20

<210> 15
 <211> 20
 <212> DNA
 <213> Mus musculus

<400> 15
 gagatgagga atatgcacgg 20

<210> 16
 <211> 18
 <212> DNA
 <213> Homo sapiens

<400> 16
 gacatctgac tcggctgc 18

<210> 17
 <211> 17
 <212> DNA
 <213> Homo sapiens

<400> 17
 tcacgcagta aaccaac 17

<210> 18
 <211> 948
 <212> PRT
 <213> Mus musculus

<400> 18
 Met Pro Ser Leu Pro Ala Pro Pro Ala Pro Arg Leu Leu Leu Gly Leu
 1 5 10 15

Leu Leu Leu Gly Ser Arg Pro Ala Ser Gly Thr Gly Pro Glu Pro Pro
 20 25 30

Ala Leu Pro Ile Arg Ser Glu Lys Glu Pro Leu Pro Val Arg Gly Ala
 35 40 45

<210> 19
 <211> 176
 <212> PRT
 <213> Rattus norvegicus

<400> 19
 Gly Gly Leu Arg Leu Ala Ser Glu Gly Val Arg Met Ser Leu Ala Pro
 1 5 10 15
 Asn Gly Glu Ala Ala Thr Ser Pro Met Leu Pro Ala Gly Pro Gly Pro
 20 25 30
 Glu Ala Pro Val Pro Ala Lys His Gly Ser Ser Gly Arg Pro Arg Asp
 35 40 45
 Pro Asn Thr Cys Phe Phe Glu Gly Gln Gln Arg Pro His Gly Ala Arg
 50 55 60
 Trp Ala Pro Asn Tyr Asp Pro Leu Cys Ser Leu Cys Thr Cys Gln Arg
 65 70 75 80
 Arg Thr Val Ile Cys Asp Pro Val Val Cys Pro Pro Pro Arg Cys Ser
 85 90 95
 Gln Pro Val Gln Ala Leu Asp Gln Trp Cys Pro Val Cys Ser Glu Lys
 100 105 110
 Gln Arg Ser Arg Asp Leu Ser Ser Leu Pro Asn Leu Glu Pro Gly Glu
 115 120 125
 Gly Cys Tyr Phe Asp Gly Asp Arg Ser Trp Arg Ala Ala Gly Thr Arg
 130 135 140
 Trp His Pro Val Val Pro Pro Phe Gly Leu Ile Lys Cys Gly Val Cys
 145 150 155 160
 Thr Cys Lys Gly Val Asn Gly Glu Val His Ser Glu Lys Val Gln Cys
 165 170 175

<210> 20
 <211> 801
 <212> PRT
 <213> Homo sapiens

<400> 20
 Gln Val Ala Ala Gly His Cys Cys Gln Thr Cys Pro Gln Glu Arg Ser
 1 5 10 15
 Ser Ser Glu Arg Gln Pro Ser Gly Leu Ser Phe Glu Tyr Pro Arg Asp
 20 25 30
 Pro Glu His Arg Ser Tyr Ser Asp Arg Gly Glu Pro Gly Ala Glu Glu

35					40					45								
Arg	Ala	Arg	Gly	Asp	Gly	His	Thr	Asp	Phe	Val	Ala	Leu	Leu	Thr	Gly			
50					55					60								
Pro	Arg	Ser	Gln	Ala	Val	Ala	Arg	Ala	Arg	Ala	Ser	Leu	Leu	Arg	Ser			
65					70					75					80			
Ser	Leu	Arg	Phe	Ser	Ile	Ser	Tyr	Arg	Arg	Leu	Asp	Arg	Pro	Thr	Arg			
					85					90					95			
Ile	Arg	Phe	Ser	Asp	Pro	Asn	Gly	Ser	Val	Leu	Phe	Glu	His	Pro	Ala			
					100					105					110			
Ala	Pro	Thr	Gln	Asp	Gly	Leu	Val	Cys	Gly	Val	Trp	Arg	Ala	Val	Pro			
					115					120					125			
Arg	Leu	Ser	Leu	Arg	Leu	Leu	Arg	Ala	Glu	Gln	Leu	His	Val	Ala	Leu			
					130					135					140			
Val	Thr	Leu	Thr	His	Pro	Ser	Gly	Glu	Val	Trp	Gly	Pro	Leu	Ile	Arg			
					145					150					155		160	
His	Arg	Ala	Leu	Ala	Ala	Glu	Thr	Phe	Ser	Ala	Ile	Leu	Thr	Leu	Glu			
					165					170					175			
Gly	Pro	Pro	Gln	Gln	Gly	Val	Gly	Gly	Ile	Thr	Leu	Leu	Thr	Leu	Ser			
					180					185					190			
Asp	Thr	Glu	Asp	Ser	Leu	His	Phe	Leu	Leu	Leu	Phe	Arg	Gly	Leu	Leu			
					195					200					205			
Glu	Pro	Arg	Ser	Gly	Gly	Leu	Thr	Gln	Val	Pro	Leu	Arg	Leu	Gln	Ile			
					210					215					220			
Leu	His	Gln	Gly	Gln	Leu	Leu	Arg	Glu	Leu	Gln	Ala	Asn	Val	Ser	Ala			
					225					230					235		240	
Gln	Glu	Pro	Gly	Phe	Ala	Glu	Val	Leu	Pro	Asn	Leu	Thr	Val	Gln	Glu			
					245					250					255			
Met	Asp	Trp	Leu	Val	Leu	Gly	Glu	Leu	Gln	Met	Ala	Leu	Glu	Trp	Ala			
					260					265					270			
Gly	Arg	Pro	Gly	Leu	Arg	Ile	Ser	Gly	His	Ile	Ala	Ala	Arg	Lys	Ser			
					275					280					285			
Cys	Asp	Val	Leu	Gln	Ser	Val	Leu	Cys	Gly	Ala	Asp	Ala	Leu	Ile	Pro			
					290					295					300			
Val	Gln	Thr	Gly	Ala	Ala	Gly	Ser	Ala	Ser	Leu	Thr	Leu	Leu	Gly	Asn			
					305					310					315		320	
Gly	Ser	Leu	Ile	Tyr	Gln	Ala	Val	Gly	Ile	Cys	Pro	Gly	Leu	Gly	Ala			
					325					330					335			
Arg	Gly	Ala	His	Met	Leu	Leu	Gln	Asn	Glu	Leu	Phe	Leu	Asn	Val	Gly			

gctagcggcc gcgccaccat gccgagcctc ccggccccg

39

<210> 24

<211> 36

<212> DNA

<213> Mus musculus

<400> 24

ggatccgtcg acggagtgtc ccgcttcttt ctccag

36

<210> 25

<211> 1341

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (1)..(1341)

<400> 25

atg gga ggc atg aaa tac atc ttt tcg ttg ttg ttc ttt ctt ttg cta 48
Met Gly Gly Met Lys Tyr Ile Phe Ser Leu Leu Phe Phe Leu Leu Leu
1 5 10 15

gaa gga ggc aaa aca gag caa gta aaa cat tca gag aca tat tgc atg 96
Glu Gly Gly Lys Thr Glu Gln Val Lys His Ser Glu Thr Tyr Cys Met
20 25 30

ttt caa gac aag aag tac aga gtg ggt gag aga tgg cat cct tac ctg 144
Phe Gln Asp Lys Lys Tyr Arg Val Gly Glu Arg Trp His Pro Tyr Leu
35 40 45

gaa cct tat ggg ttg gtt tac tgc gtg aac tgc atc tgc tca gag aat 192
Glu Pro Tyr Gly Leu Val Tyr Cys Val Asn Cys Ile Cys Ser Glu Asn
50 55 60

ggg aat gtg ctt tgc agc cga gtc aga tgt cca aat gtt cat tgc ctt 240
Gly Asn Val Leu Cys Ser Arg Val Arg Cys Pro Asn Val His Cys Leu
65 70 75 80

tct cct gtg cat att cct cat ctg tgc tgc cct cgc tgc cca gaa gac 288
Ser Pro Val His Ile Pro His Leu Cys Cys Pro Arg Cys Pro Glu Asp
85 90 95

tcc tta ccc cca gtg aac aat aag gtg acc agc aag tct tgc gag tac 336
Ser Leu Pro Pro Val Asn Asn Lys Val Thr Ser Lys Ser Cys Glu Tyr
100 105 110

aat ggg aca act tac caa cat gga gag ctg ttc gta gct gaa ggg ctc 384
Asn Gly Thr Thr Tyr Gln His Gly Glu Leu Phe Val Ala Glu Gly Leu
115 120 125

ttt cag aat cgg caa ccc aat caa tgc acc cag tgc agc tgt tcg gag 432
Phe Gln Asn Arg Gln Pro Asn Gln Cys Thr Gln Cys Ser Cys Ser Glu
130 135 140

gga aac gtg tat tgt ggt ctc aag act tgc ccc aaa tta acc tgt gcc	480
Gly Asn Val Tyr Cys Gly Leu Lys Thr Cys Pro Lys Leu Thr Cys Ala	
145 150 155 160	
ttc cca gtc tct gtt cca gat tcc tgc tgc cgg gta tgc aga gga gat	528
Phe Pro Val Ser Val Pro Asp Ser Cys Cys Arg Val Cys Arg Gly Asp	
165 170 175	
gga gaa ctg tca tgg gaa cat tct gat ggt gat atc ttc cgg caa cct	576
Gly Glu Leu Ser Trp Glu His Ser Asp Gly Asp Ile Phe Arg Gln Pro	
180 185 190	
gcc aac aga gaa gca aga cat tct tac cac cgc tct cac tat gat cct	624
Ala Asn Arg Glu Ala Arg His Ser Tyr His Arg Ser His Tyr Asp Pro	
195 200 205	
cca cca agc cga cag gct gga ggt ctg tcc cgc ttt cct ggg gcc aga	672
Pro Pro Ser Arg Gln Ala Gly Gly Leu Ser Arg Phe Pro Gly Ala Arg	
210 215 220	
agt cac cgg gga gct ctt atg gat tcc cag caa gca tca gga acc att	720
Ser His Arg Gly Ala Leu Met Asp Ser Gln Gln Ala Ser Gly Thr Ile	
225 230 235 240	
gtg caa att gtc atc aat aac aaa cac aag cat gga caa gtg tgt gtt	768
Val Gln Ile Val Ile Asn Asn Lys His Lys His Gly Gln Val Cys Val	
245 250 255	
tcc aat gga aag acc tat tct cat ggc gag tcc tgg cac cca aac ctc	816
Ser Asn Gly Lys Thr Tyr Ser His Gly Glu Ser Trp His Pro Asn Leu	
260 265 270	
cgg gca ttt ggc att gtg gag tgt gtg cta tgt act tgt aat gtc acc	864
Arg Ala Phe Gly Ile Val Glu Cys Val Leu Cys Thr Cys Asn Val Thr	
275 280 285	
aag caa gag tgt aag aaa atc cac tgc ccc aat cga tac ccc tgc aag	912
Lys Gln Glu Cys Lys Lys Ile His Cys Pro Asn Arg Tyr Pro Cys Lys	
290 295 300	
tat cct caa aaa ata gac gga aag tgc tgc aag gtg tgt cca gaa gaa	960
Tyr Pro Gln Lys Ile Asp Gly Lys Cys Cys Lys Val Cys Pro Glu Glu	
305 310 315 320	
ctt cca ggc caa agc ttt gac aat aaa ggc tac ttc tgc ggg gaa gaa	1008
Leu Pro Gly Gln Ser Phe Asp Asn Lys Gly Tyr Phe Cys Gly Glu Glu	
325 330 335	
acg atg cct gtg tat gag tct gta ttc atg gag gat ggg gag aca acc	1056
Thr Met Pro Val Tyr Glu Ser Val Phe Met Glu Asp Gly Glu Thr Thr	
340 345 350	
aga aaa ata gca ctg gag act gag aga cca cct cag gta gag gtc cac	1104
Arg Lys Ile Ala Leu Glu Thr Glu Arg Pro Pro Gln Val Glu Val His	
355 360 365	

gtt tgg act att cga aag ggc att ctc cag cac ttc cat att gag aag	1152
Val Trp Thr Ile Arg Lys Gly Ile Leu Gln His Phe His Ile Glu Lys	
370 375 380	
atc tcc aag agg atg ttt gag gag ctt cct cac ttc aag ctg gtg acc	1200
Ile Ser Lys Arg Met Phe Glu Glu Leu Pro His Phe Lys Leu Val Thr	
385 390 395 400	
aga aca acc ctg agc cag tgg aag atc ttc acc gaa gga gaa gct cag	1248
Arg Thr Thr Leu Ser Gln Trp Lys Ile Phe Thr Glu Gly Glu Ala Gln	
405 410 415	
atc agc cag atg tgt tca agt cgt gta tgc aga aca gag ctt gaa gat	1296
Ile Ser Gln Met Cys Ser Ser Arg Val Cys Arg Thr Glu Leu Glu Asp	
420 425 430	
tta gtc aag gtt ttg tac ctg gag aga tct gaa aag ggc cac tgt	1341
Leu Val Lys Val Leu Tyr Leu Glu Arg Ser Glu Lys Gly His Cys	
435 440 445	

<210> 26
 <211> 447
 <212> PRT
 <213> Homo sapiens

<400> 26
Met Gly Gly Met Lys Tyr Ile Phe Ser Leu Leu Phe Phe Leu Leu Leu
1 5 10 15
Glu Gly Gly Lys Thr Glu Gln Val Lys His Ser Glu Thr Tyr Cys Met
20 25 30
Phe Gln Asp Lys Lys Tyr Arg Val Gly Glu Arg Trp His Pro Tyr Leu
35 40 45
Glu Pro Tyr Gly Leu Val Tyr Cys Val Asn Cys Ile Cys Ser Glu Asn
50 55 60
Gly Asn Val Leu Cys Ser Arg Val Arg Cys Pro Asn Val His Cys Leu
65 70 75 80
Ser Pro Val His Ile Pro His Leu Cys Cys Pro Arg Cys Pro Glu Asp
85 90 95
Ser Leu Pro Pro Val Asn Asn Lys Val Thr Ser Lys Ser Cys Glu Tyr
100 105 110
Asn Gly Thr Thr Tyr Gln His Gly Glu Leu Phe Val Ala Glu Gly Leu
115 120 125
Phe Gln Asn Arg Gln Pro Asn Gln Cys Thr Gln Cys Ser Cys Ser Glu
130 135 140
Gly Asn Val Tyr Cys Gly Leu Lys Thr Cys Pro Lys Leu Thr Cys Ala
145 150 155 160

<212> PRT

<213> Homo sapiens

<400> 27

Glu Gln Val Lys His Ser Glu Thr Tyr Cys Met Phe Gln Asp Lys Lys
1 5 10 15
Tyr Arg Val Gly Glu Arg Trp His Pro Tyr Leu Glu Pro Tyr Gly Leu
20 25 30
Val Tyr Cys Val Asn Cys Ile Cys Ser Glu Asn Gly Asn Val Leu Cys
35 40 45
Ser Arg Val Arg Cys Pro Asn Val His Cys Leu Ser Pro Val His Ile
50 55 60
Pro His Leu Cys Cys Pro Arg Cys Pro Glu Asp Ser Leu Pro Pro Val
65 70 75 80
Asn Asn Lys Val Thr Ser Lys Ser Cys Glu Tyr Asn Gly Thr Thr Tyr
85 90 95
Gln His Gly Glu Leu Phe Val Ala Glu Gly Leu Phe Gln Asn Arg Gln
100 105 110
Pro Asn Gln Cys Thr Gln Cys Ser Cys Ser Glu Gly Asn Val Tyr Cys
115 120 125
Gly Leu Lys Thr Cys Pro Lys Leu Thr Cys Ala Phe Pro Val Ser Val
130 135 140
Pro Asp Ser Cys Cys Arg Val Cys Arg Gly Asp Gly Glu Leu Ser Trp
145 150 155 160
Glu His Ser Asp Gly Asp Ile Phe Arg Gln Pro Ala Asn Arg Glu Ala
165 170 175
Arg His Ser Tyr His Arg Ser His Tyr Asp Pro Pro Pro Ser Arg Gln
180 185 190
Ala Gly Gly Leu Ser Arg Phe Pro Gly Ala Arg Ser His Arg Gly Ala
195 200 205
Leu Met Asp Ser Gln Gln Ala Ser Gly Thr Ile Val Gln Ile Val Ile
210 215 220
Asn Asn Lys His Lys His Gly Gln Val Cys Val Ser Asn Gly Lys Thr
225 230 235 240
Tyr Ser His Gly Glu Ser Trp His Pro Asn Leu Arg Ala Phe Gly Ile
245 250 255
Val Glu Cys Val Leu Cys Thr Cys Asn Val Thr Lys Gln Glu Cys Lys
260 265 270
Lys Ile His Cys Pro Asn Arg Tyr Pro Cys Lys Tyr Pro Gln Lys Ile
275 280 285

115					120					125						
Val	Tyr	Cys	Gly	Leu	Lys	Thr	Cys	Pro	Lys	Leu	Thr	Cys	Ala	Phe	Pro	
130					135					140						
Val	Ser	Val	Pro	Asp	Ser	Cys	Cys	Arg	Val	Cys	Arg	Gly	Asp	Gly	Glu	
145					150					155					160	
Leu	Ser	Trp	Glu	His	Ser	Asp	Gly	Asp	Ile	Phe	Arg	Gln	Pro	Ala	Asn	
165					170					175						
Arg	Glu	Ala	Arg	His	Ser	Tyr	His	Arg	Ser	His	Tyr	Asp	Pro	Pro	Pro	
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210					215					220						
Ile	Val	Ile	Asn	Asn	Lys	His	Lys	His	Gly	Gln	Val	Cys	Val	Ser	Asn	
225					230					235					240	
Gly	Lys	Thr	Tyr	Ser	His	Gly	Glu	Ser	Trp	His	Pro	Asn	Leu	Arg	Ala	
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Phe	Gly	Ile	Val	Glu	Cys	Val	Leu	Cys	Thr	Cys	Asn	Val	Thr	Lys	Gln	
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Glu	Cys	Lys	Lys	Ile	His	Cys	Pro	Asn	Arg	Tyr	Pro	Cys	Lys	Tyr	Pro	
275					280					285						
Gln	Lys	Ile	Asp	Gly	Lys	Cys	Cys	Lys	Val	Cys	Pro	Glu	Glu	Leu	Pro	
290					295					300						
Gly	Gln	Ser	Phe	Asp	Asn	Lys	Gly	Tyr	Phe	Cys	Gly	Glu	Glu	Thr	Met	
305					310					315					320	
Pro	Val	Tyr	Glu	Ser	Val	Phe	Met	Glu	Asp	Gly	Glu	Thr	Thr	Arg	Lys	
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Ile	Ala	Leu	Glu	Thr	Glu	Arg	Pro	Pro	Gln	Val	Glu	Val	His	Val	Trp	
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Thr	Ile	Arg	Lys	Gly	Ile	Leu	Gln	His	Phe	His	Ile	Glu	Lys	Ile	Ser	
355					360					365						
Lys	Arg	Met	Phe	Glu	Glu	Leu	Pro	His	Phe	Lys	Leu	Val	Thr	Arg	Thr	
370					375					380						
Thr	Leu	Ser	Gln	Trp	Lys	Ile	Phe	Thr	Glu	Gly	Glu	Ala	Gln	Ile	Ser	
385					390					395					400	
Gln	Met	Cys	Ser	Ser	Arg	Val	Cys	Arg	Thr	Glu	Leu	Glu	Asp	Leu	Val	
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38

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33

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 35 40 45
 Met Arg Cys Val Leu Cys Ala Cys Glu Ala Pro Gln Trp Ala Arg Arg
 50 55 60
 Gly Arg Gly Pro Gly Arg Val Ser Cys Lys Asn Ile Lys Pro Gln Cys
 65 70 75 80
 Pro Thr Leu Ala Cys Arg Gln Pro Arg Gln Leu Pro Gly His Cys Cys
 85 90 95
 Gln Thr Cys Pro Gln Glu Arg Ser Asn Leu Asp Pro Gln Pro Ala Gly
 100 105 110
 Leu Val Phe Glu Tyr Pro Arg Asp Pro Glu His Arg Ser Tyr Ser Asp
 115 120 125
 Arg Gly Glu Pro Gly Val Gly Glu Arg Thr Arg Ala Asp Gly His Thr
 130 135 140
 Asp Phe Val Ala Leu Leu Thr Gly Pro Arg Ser Gln Ala Val Ala Arg
 145 150 155 160

Ala Arg Val Ser Leu Leu Arg Ser Ser Leu Arg Phe Ser Val Ser Tyr
165 170 175

Gln Arg Leu Asp Arg Pro Ser Arg Val Arg Phe Thr Asp Pro Thr Gly
180 185 190

Asn Ile Leu Phe Glu His Pro Ala Thr Pro Thr Gln Asp Gly Leu Val
195 200 205

Cys Gly Val Trp Arg Ala Val Pro Arg Leu Ser Val Arg Leu Leu Arg
210 215 220

Ala Glu Gln Leu Arg Val Ala Leu Val Thr Ser Thr His Pro Ser Gly
225 230 235 240

Glu Val Trp Gly Pro Leu Ile Trp Gln Gly Ala Leu Ala Ala Glu Thr
245 250 255

Phe Ser Ala Ile Leu Thr Leu Glu Asp Pro Leu Gln Arg Gly Val Gly
260 265 270

Gly Ile Ala Leu Leu Thr Leu Ser Asp Thr Glu Asp Ser Leu His Phe
275 280 285

Leu Leu Leu Phe Arg Gly Leu Leu Gly Gly Leu Ala Gln Ala Pro Leu
290 295 300

Lys Leu Gln Ile Leu His Gln Gly Gln Leu Leu Arg Glu Leu Gln Ala
305 310 315 320

Asn Thr Ser Ala Gln Glu Pro Gly Phe Ala Glu Val Leu Pro Ser Leu
325 330 335

Thr Asp Gln Glu Met Asp Trp Leu Glu Leu Gly Glu Leu Gln Met Val
340 345 350

Leu Glu Lys Ala Gly Gly Pro Glu Leu Arg Ile Ser Gly Tyr Ile Thr
355 360 365

Thr Arg Gln Ser Cys Asp Val Leu Gln Ser Val Leu Cys Gly Ala Asp
370 375 380

Ala Leu Ile Pro Val Gln Thr Gly Ala Ala Gly Ser Ala Ser Phe Ile
385 390 395 400

Leu Leu Gly Asn Gly Ser Leu Ile Tyr Gln Val Gln Val Val Gly Thr
405 410 415

Gly Ser Glu Val Val Ala Met Thr Leu Glu Thr Lys Pro Gln Arg Lys
420 425 430

Asn Gln Arg Thr Val Leu Cys His Met Ala Gly Leu Gln Pro Gly Gly
435 440 445

His Met Ala Val Gly Met Cys Ser Gly Leu Gly Ala Arg Gly Ala His
450 455 460

SECRET 5164260

Sub
a1

Met Leu Leu Gln Asn Glu Leu Phe Leu Asn Val Gly Thr Lys Asp Phe
465 470 475 480
Pro Asp Gly Glu Leu Arg Gly His Val Thr Ala Leu Cys Tyr Ser Gly
485 490 495
His Ser Ala Arg Tyr Asp Arg Leu Pro Val Pro Leu Ala Gly Ala Leu
500 505 510
Val Leu Pro Pro Val Arg Ser Gln Ala Ala Gly His Ala Trp Leu Ser
515 520 525
Leu Asp Thr His Cys His Leu His Tyr Glu Val Leu Leu Ala Gly Leu
530 535 540
Gly Gly Ser Glu Gln Gly Thr Val Thr Ala His Leu Leu Gly Pro Pro
545 550 555 560
Gly Met Pro Gly Pro Gln Arg Leu Leu Lys Gly Phe Tyr Gly Ser Glu
565 570 575
Ala Gln Gly Val Val Lys Asp Leu Glu Pro Val Leu Leu Arg His Leu
580 585 590
Ala Gln Gly Thr Ala Ser Leu Leu Ile Thr Thr Lys Ser Ser Pro Arg
595 600 605
Gly Glu Leu Arg Gly Gln Val His Ile Ala Ser Gln Cys Glu Ala Gly
610 615 620
Gly Leu Arg Leu Ala Ser Glu Gly Val Gln Met Pro Leu Ala Pro Asn
625 630 635 640
Gly Glu Ala Ala Thr Ser Pro Met Leu Pro Ala Gly Pro Gly Pro Glu
645 650 655
Ala Pro Val Pro Ala Lys His Gly Ser Pro Gly Arg Pro Arg Asp Pro
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Asn Thr Cys Phe Phe Glu Gly Gln Gln Arg Pro His Gly Ala Arg Trp
675 680 685
Ala Pro Asn Tyr Asp Pro Leu Cys Ser Leu Cys Ile Cys Gln Arg Arg
690 695 700
Thr Val Ile Cys Asp Pro Val Val Cys Pro Pro Pro Ser Cys Pro His
705 710 715 720
Pro Val Gln Ala Leu Asp Gln Cys Cys Pro Val Cys Pro Glu Lys Gln
725 730 735
Arg Ser Arg Asp Leu Pro Ser Leu Pro Asn Leu Glu Pro Gly Glu Gly
740 745 750
Cys Tyr Phe Asp Gly Asp Arg Ser Trp Arg Ala Ala Gly Thr Arg Trp
755 760 765

His	Pro	Val	Val	Pro	Pro	Phe	Gly	Leu	Ile	Lys	Cys	Ala	Val	Cys	Thr	
770						775					780					
Cys	Lys	Gly	Ala	Thr	Gly	Glu	Val	His	Cys	Glu	Lys	Val	Gln	Cys	Pro	
785					790					795					800	
Arg	Leu	Ala	Cys	Ala	Gln	Pro	Val	Arg	Ala	Asn	Pro	Thr	Asp	Cys	Cys	
			805						810					815		
Lys	Gln	Cys	Pro	Val	Gly	Ser	Gly	Thr	Asn	Ala	Lys	Leu	Gly	Asp	Pro	
			820					825					830			
Met	Gln	Ala	Asp	Gly	Pro	Arg	Gly	Cys	Arg	Phe	Ala	Gly	Gln	Trp	Phe	
		835					840					845				
Pro	Glu	Asn	Gln	Ser	Trp	His	Pro	Ser	Val	Pro	Pro	Phe	Gly	Glu	Met	
	850					855					860					
Ser	Cys	Ile	Thr	Cys	Arg	Cys	Gly	Ala	Gly	Val	Pro	His	Cys	Glu	Arg	
865					870					875					880	
Asp	Asp	Cys	Ser	Pro	Pro	Leu	Ser	Cys	Gly	Ser	Gly	Lys	Glu	Ser	Arg	
				885					890					895		
Cys	Cys	Ser	His	Cys	Thr	Ala	Gln	Arg	Ser	Ser	Glu	Thr	Arg	Thr	Leu	
			900					905					910			
Pro	Glu	Leu	Glu	Lys	Glu	Ala	Glu	His	Ser	Val	Asp	Gly	Ser	Gly	Thr	
		915					920					925				
Asp	Tyr	Lys	Asp	Asp	Asp	Asp	Lys									
	930					935										